Amendments to the claims:

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of claims:

1-26. (Canceled)

27. (Currently amended) An isolated peptide which binds to a DM2 protein, which peptide comprises an amino acid motif comprising at least the eight consecutive amino acids from F to R₄ of the formula

$$R_1-X-F-X-R_2-R_3-W-X-X-R_4(I)$$

(SEQ ID NO: 4)

wherein

R₁ is a proline (P), leucine (L), glutamic acid (E), cysteine (C) or glutamine (Q),

X stands for any natural amino acid,

R₂ is arginine (R), histidine (H), **glutamic acid (E)**, **cysteine (C)**, **serine** (S), or aspartic acid (D),

R₃ is histidine (H), phenylalanine (F) or tyrosine (Y),

R₄ is phenylalanine (F), glutamine (Q) or leucine (L); and

F is phenylalanine and W is tryptophan,

and inhibits the binding of said DM2 protein to a p53 protein.

28. (Previously presented) The peptide according to claim 27 wherein the peptide binds to human DM2 (HDM2).

- 29. (Previously presented) The peptide according to claim 27, which is coupled to a biotin moiety.
- 30. (Previously presented) The peptide according to claim 27, which is a cyclic peptide.
- 31. (Previously presented) The peptide according to claim 27, which is a cyclic lactam.
- 32. (Previously presented) The peptide according to claim 27 which comprises a disulfide bond.
- 33. (Previously presented) The peptide according to claim 27 which comprises no more than fifteen amino acids (15 mers).
- 34. (Previously presented) The peptide according to claim 27 which comprises an amino acid motif selected from the group consisting of M-P-R-F-M-D-Y-W-E-G-L-N (SEQ ID NO: 6), Q-P-T-F-S-D-Y-W-K-L-L-P (SEQ ID NO: 7), and P-X-F-X-D-Y-W-X-X-L (SEQ ID NO: 8).
- 35. (Currently amended) An isolated peptide which comprises eight amino acids according to the formula

$$F-X_2-R_2-R_3-W-X_3-X_4-R_4$$
 (Ib) (SEQ ID NO: 10)

wherein R_2 is arginine (R), histidine (H), glutamic acid (E), cysteine (C), serine (S), or aspartic acid (D);

R₃ is histidine (H), phenylalanine (F), or tyrosine (Y);

R₄ is phenylalanine (F), glutamine (Q) or leucine (L);

X₂ is methionine (M), isoleucine (I), threonine (T), arginine (R), alanine (A) or serine (S);

X₃ is glutamic acid (E), threonine (T), alanine (A), phenylalanine (F) or serine (S); and

X₄ is glycine (G), glutamine (Q), threonine (T), alanine (A) or aspartic acid (D).

36. (Currently amended) The peptide according to claim 35 comprising an amino acid motif of the formula

$$X_1$$
-F- X_2 -R₂-R₃- W - X_3 - X_4 -R₄ (Ic) (SEQ ID NO: 11)

wherein

R₂ is arginine (R), histidine (H), glutamic acid (E), cysteine (C), serine (S), or aspartic acid (D);

R₃ is histidine (H), phenylalanine (F) or tyrosine (Y);

R₄ is phenylalanine (F), glutamine (Q) or leucine (L);

X₁ is arginine (R), asparagine (N), alanine (A), threonine (T), or valine (V);

X₂ is methionine (M), isoleucine (I), threonine (T), arginine (R), alanine (A), or serine (S);

X₃ is glutamic acid (E), threonine (T), alanine (A), phenylalanine (F), or serine (S); and

X₄ is glycine (G), glutamine (Q), threonine (T), alanine (A), or aspartic acid (D).

37. (Canceled)

38. (Previously presented) The peptide according to claim 27, wherein R₂ is aspartic acid (D).

39. (Previously presented) The peptide according to claim 35, wherein at least one of R_2 , X_2 , X_3 , and X_4 is defined as follows: R_2 is aspartic acid (D), X_2 is methionine (M), X_3 is glutamic acid (E), and X_4 is glycine (G).

40. (Previously presented) The peptide according to claim 36, wherein at least one of R_2 , X_1 , X_2 , X_3 , and X_4 is defined as follows: R_2 is aspartic acid (D), X_1 is arginine (R), X_2 is methionine (M), X_3 is glutamic acid (E), and X_4 is glycine (G).

41. (Currently amended) A method for inhibiting the <u>in vitro</u> binding of a DM2 protein to a p53 protein comprising contacting said DM2 protein with a peptide <u>in vitro</u>, which peptide comprises an amino acid motif comprising at least eight consecutive amino acids of the formula

$$R_1-X-F-X-R_2-R_3-W-X-X-R_4(I)$$
 (SEQ ID NO: 4)

wherein

R₁ is a proline (P), leucine (L), glutamic acid (E), cysteine (C) or glutamine (Q),

X stands for any natural amino acid,

R₂ is arginine (R), histidine (H), **glutamic acid (E)**, **cysteine (C)**, **serine** (S), or aspartic acid (D),

R₃ is histidine (H), phenylalanine (F) or tyrosine (Y),

R₄ is phenylalanine (F), glutamine (Q) or leucine (L); and

F is phenylalanine and W is tryptophan,

and inhibits the binding of said DM2 protein to a p53 protein.

42. (Previously presented) The method of claim 41 wherein R_2 is aspartic acid (D).

43-51. (Canceled)

52. (Currently amended) A composition comprising an isolated peptide, which peptide comprises an amino acid motif comprising at least eight consecutive amino acids of the formula

$$R_1$$
-X-F- X - R_2 - R_3 - W -X-X- R_4 (I) (SEQ ID NO: 4)

wherein

R₁ is a proline (P), leucine (L), glutamic acid (E), cysteine (C) or glutamine (Q),

X stands for any natural amino acid,

R₂ is arginine (R), histidine (H), glutamic acid (E), cysteine (C), serine (S), or aspartic acid (D),

R₃ is histidine (H), phenylalanine (F) or tyrosine (Y),

R₄ is phenylalanine (F), glutamine (Q) or leucine (L); and

F is phenylalanine and W is tryptophan,

and inhibits the binding of said DM2 protein to a p53 protein, in admixture with at least one pharmaceutically acceptable carrier.